

## SEQUENCE LISTING

<110> Salceda, Susana  
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 Sun, Yongming

<120> Compositions and Methods Relating to Breast Specific Genes and Proteins

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<150> 60/268,289

<151> 2001-02-13

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703

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 gggagaatgg gttaccgggt cacagttcca cacatttgcg agacaacaga cgggagaaga 600  
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 <213> Homo sapien

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 <211> 337  
 <212> DNA  
 <213> Homo sapien

<400> 29  
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 atttagatgt ttttaacata atggcaaact aaaatgt 337

<210> 30  
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 <212> DNA  
 <213> Homo sapien

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 <211> 260  
 <212> DNA  
 <213> Homo sapien

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 attaattaaa aagactttta gacaacctct taaatggaat tacactatgg aaaacagggc 180  
 tcccccaaaa acacctaggc agaactgaga gttctttgaa aaccattccc aataaaaact 240  
 aaatgaaaaa taaatataaa 260

<210> 32  
 <211> 1416  
 <212> DNA  
 <213> Homo sapien

<400> 32  
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 tttaggggaag ttgttgacat ccaaatcaca gaaccaaggt caaaagcaaa atacaaaggt 180  
 accctcaaaa atattttaca tgaagtaaat acactaacag aattttaaac aggtacaaaa 240  
 tattgaaatg accaacgtta catgatttca agggttgtcc tttctgtgct ttttatctgt 300  
 cacgacagga aggtgtggaa agtttatatc cttaatttga ctactcttgg atattaaaat 360  
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 cagggctccc tcaaaaacac ctaggcagaa ctgagagttc tttgaaaacc attcccaata 480  
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<212> DNA
<213> Homo sapien
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<211> 1344
<212> DNA
<213> Homo sapien
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gcttttagca gagagaagcc tgtatatgtt acatgtgtga ctttcagtag tttaaaqaqa      240
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<210> 35  
 <211> 163  
 <212> DNA  
 <213> Homo sapien

<400> 35	
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ttctaaaccc aaaagtgcta cctaagaaga aatttagcca aaaaataccc agctaaggta	120
gccatagcca agtgtattta agtatgttat agaatatatt tga	163

<210> 36  
 <211> 643  
 <212> DNA  
 <213> Homo sapien

<400> 36	
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 caagagaagt gataatggat gataatggaa ttgatactgt atttaggatc ctttggttgt 240  
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 gaatatattt gaaagcttcc ttccagtttg agctttgtat ctgctgtgga actgttatgg 480  
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 aacaagaata ccaaatagaa tacgaaataa taaagataaa ccaaagaat accaaataat 600  
 aaagattttt aagaaatgga aaaaaaaaaa aaaaaaaaaa att 643

<210> 37  
 <211> 478  
 <212> DNA  
 <213> Homo sapien

<400> 37  
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 gctcattacg cggtttccct ggtggtggac attgggtttc tccgctccac aattccccag 420  
 acaacttagg gacgcaagaa accccgatca caaaagcact ccacaacca cacacaca 478

<210> 38  
 <211> 833  
 <212> DNA  
 <213> Homo sapien

<400> 38  
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 ccctagtctt ctataaaaga taatccactt tatcgtact acgattccgt tatttataga 180  
 aagagaagat cgttctcgta gtacacatgt ttatggagga atatcttaag atagaacact 240

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<212> DNA  
<213> Homo sapien

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tcacagagaa ttttaatgac attggaaaat gtaagaaatt tgaaaaaaag atggagtaaa 180  
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cactcccaa cagttaaaac cagctcta atccaatctg cagagtttta agcaaagcc 660  
ggattgtctg gacagagaaa atcctccaga ggagagccag agaaaataga tgtgaggg 718

<210> 40  
<211> 1439  
<212> DNA  
<213> Homo sapien

<400> 40  
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 caaccagtca aaattcaaca tctttaagaa tattgtact ttgggcaaaa tttgagtttc 1380  
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<210> 41  
 <211> 298  
 <212> DNA  
 <213> Homo sapien

<400> 41  
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aagtgctgga aaattagggc aggaattacg tgtttgcaag ttgtgccatc actgggttga 240  
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<210> 42  
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 <212> DNA  
 <213> Homo sapien

<400> 42  
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<210> 43  
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 <212> DNA  
 <213> Homo sapien

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<220>  
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 <222> (262)..(262)  
 <223> a, c, g or t

<400> 44  
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 tctaaaagaa aaaaagaaaa agaaatagaa catttccaga tctcagaagt cttctcttgt 120  
 cactatccct tacaaaggca acctgacttt taataccata gattaatttt gtctgttttt 180  
 atactttata taaatgtaat caatcaatat gcaatctttn gtgtcagctt cttntgtctt 240  
 acattatact tgtgagatcc anaaaaaaaa aaaaaacaaa aaaaaaaaaa acggcttggg 300  
 gcggtaacct caaggcggcc aataaggcgg ggtctcgcg gtggtggaaa tatgggtgta 360  
 tactcgggcg ctcaaaatat cccaacacac aactatat caagcggcac ggcaaaaaag 420  
 ggggaaaacc gaaaacaaga aaacagaaaa aaaaaaagaa aaaaaaaaaa aaacagaaaa 480  
 aaaaaaaaaa acgaa 495

<210> 45  
 <211> 651  
 <212> DNA  
 <213> Homo sapien

<400> 45  
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 tccacattgt tgccaacatt cttaatcttg tgttttttaa taacagctat cctaacaggt 120  
 atgaggtgat ctctctcatt gcggttttga ttgcatttc cctaacggtt ggtgatactg 180  
 agcatttttg catacaccgg gtcatttgtt ctttgttgtt gacttgagat cccttatata 240  
 gtttggatac tgctgtggcc tgaatgtttg tgtccccc aaattcgtat attgaactct 300  
 catccctaag gtcaacagtt tagggaagcg attaggtcct gaggactctg ccctcttgca 360  
 tagaattagt gctcttataa aagatgcccg agggagctct tttgcccctc ctgccatgtg 420  
 aggacacagc tagaagctac catctgtgaa ccaggaagcc cccctcacca gacactgaat 480

ctgctggagc caccatcttg gacttcccag cctccagagc tgtgagaaat acatgcctgt 540  
 agttaagcaa aaaaaaaaaa aaaacaacaa aaacagcgtg ggggaaacaa ggacaaaaga 600  
 ggtcacctgg gtaaaaggga actcggacca cattccaaca cttacacaaa g 651

<210> 46  
 <211> 873  
 <212> DNA  
 <213> Homo sapien

<400> 46  
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 ctgattccca ctgagttacg ccgagaggct ctggccttac aggggtccct ggagtttgat 180  
 gatgctggag gtgaagggtg gaccagccac gtggatgatg aataccgatg ggcaggagtc 240  
 gaggatccca aggttatgat cactacctcc cgagacccca gttcccgctt caagatgttt 300  
 gcaaaggagc tgaagctggt gttcccgggc gccagcgaa tgaaccgagg tcgacatgaa 360  
 gtgggggcac tgggtgcgagc ctgcaaagcc aacggcgctc ccgatctgct ggtcgttcac 420  
 gagcatcggg gcacacctgt ggggctcatc gtcagccacc tgccctttgg tcctactgcc 480  
 tacttcacgc tgtgcaatgt ggtcatgcgg catgacatcc cagacctggg caccatgtcg 540  
 gaggccaaag cccacctcat cacacacggc ttctctctcc gcctgggcaa gcgggtctct 600  
 gacatcctcc gatacctatt tcccggtgcc aaagatgaca gccaccgggt catcaccttc 660  
 gcaaaccagg acgactacat atcattccgg caccatgtgt ataagaagac agaccaccgc 720  
 aacgtggagc tctactgagg cgggccccgc tttgagctga agctgtacat gatccgtctg 780  
 ggcacgctgg agcaggaggc cacagcagac gtggagtggc gctggcacc ttacaccaat 840  
 accgcacgca agagagtctt cctgagcacc gag 873

<210> 47  
 <211> 213  
 <212> DNA  
 <213> Homo sapien

<400> 47  
 tatgagtata agggcatggt ttcctctaag ctgtcgagcg gcgcatgtga tggatccggg 60  
 caggctactgg acacctggca tgctgactgc cacgtgcagg caagaaacat ctgtccagta 120  
 agttaggggg aagacgggat ggggaataaa ccctcggaaa tctctgcaca ccactcttgg 180  
 tgctatgctt ttaattctgt ttccctttct cct 213



<210> 48  
 <211> 658  
 <212> DNA  
 <213> Homo sapien

<400> 48  
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 aggggaaggag acgattggag tcaactcaat gtgctcaaaa aaagaagagt cggggacctc 120  
 ctagccagtt acattccaga ggatgaggcg ctgatgcttc gggatggacg ctttgcttgt 180  
 gccatctgcc cccatcgacc ggtactggac accctggcca tgctgactgc ccaccgtgca 240  
 ggcaagaaac atctgtccag taagttaggg ggaagacggg atggggaata aaccctcgaa 300  
 atctctgcac accactcttg gtgctatgct ttttaattctg tttcccttc tctcagget 360  
 tgcagctttt ctatggcaag aagcagccgg gaaaggaaag aaagcagaat ccaaaacatc 420  
 agaatgaatt gagaaggga gaaaccaaag ctgaggctcc tctgctaact cagacacgac 480  
 ttatcaccca gagtgtctg cacagagctc cccactataa cagttgtctgc cgccggaagt 540  
 acaggtatgg gacgggaaag ccagaggtag gaaggctcag aaggagacag atggctctaa 600  
 aagagttttc cagtgtgtat tctgaggaat actagtgttc tggagatgtt acttagtg 658

<210> 49  
 <211> 703  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (169)..(169)  
 <223> a, c, g or t

<400> 49  
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 cccctggcaa cactccctt aagtgaagag tgacaacttt cctgggcant gtgctttcag 180  
 tagtatgtgg ctttacctgt ttccattaga atttttaaca ccaaattcaa gcagtgagct 240  
 tgtaactatt ctgagattat gaaatatcct tttatataca actatTTTTTg tctcaaacat 300  
 gtttctttat acataaaaaa tagatatTTTc tgTTTccatt ttttaataaa attctgtcct 360  
 tatttcagaa gtgagaaaaa tcaatactcc aatattaaaa agcaggaata accatagtTC 420  
 tattattaac tgtgggccac cacactctct gtcctactgc ttccacaga atctgaggTg 480  
 ccaagggtcg caaggccttt gagggcaagc tgcacatttt acagatgaag aaacagatcc 540

gacatgggct tgtgacatgt ccaaggtcac aaggccagtt aacagcaagc taggatgaga 600  
 atccttctta ctagaactta gtattaatat taatgcgaca gctgggtatc atgtcatagc 660  
 tgttccggtg aatgtatcgt caaaaaaaaa aaaaaaaaaa aaa 703

<210> 50  
 <211> 1251  
 <212> DNA  
 <213> Homo sapien

<400> 50  
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 atgcagttta cataattgaa atgtgttttt ctctgtgtgc tgttctcata ttccaatatt 120  
 cttttttcct ctcatgggtca tgatgttttc ttttgagata taattcacat accataaaat 180  
 tgatgctttt aaactataca attcgttagc tgggtgtggc agcacacacc tgtagtccca 240  
 gctactcagg aggctgaagt gagaggatca cttgaactgg gaggcagagg ttgcagtgaa 300  
 ccgagattgc gccgctgcac tccatcctag gcgacagggt gagcccctgt ctcaaaaata 360  
 aataaataaa caattcagtg gttcctagta cattcaaaca gttatacaac tatcaccact 420  
 attccaattc cagaacattc tcatcatcgc ccaaagaaac cacataccta ttagcagtc 480  
 ctccccatcc tccctttctc agcccctggc aaccactccc ttaagtgaag agtgacaact 540  
 ttcttgggca ttgtgctttc agtagtatgt ggctttacat gtttccatta gaatttttaa 600  
 caccaaattc aagcagtgag ctttgtaact attctgagat tatgaaatat ctttttatat 660  
 acaactatth ttgtctcaaa catgtttctt tatacataaa aaatagatat ttctgtttcc 720  
 attttttaat caaattctgt ctttatthca gaagtgaaga aaatcaatac tccaatatta 780  
 aaaagcagga ataaccatag ttctattatt aactgtgggc caccacactc tctgtcctac 840  
 tgcttccac agaattctgag gtgccaaggc ctgcaaggcc tttgagggca agctgcacat 900  
 tttacagatg aagaaacaga tccgacatgg gcttgtgaca tgtccaaggc cacaaggcca 960  
 gttaacagca aagctaggat gagaatccct tcttactaga acttttagtat caaatattta 1020  
 aatgctgact ttgtgggtaa cctaattcag ctaccacatg aatctaatta tgtcagtttc 1080  
 ctctacagct ttgatctgag catgtgattt cttttttttt accattttta aaacatttac 1140  
 atgttatctt ttaagacctg taaggacatg actagtctat ttagccagag ggcccaaadc 1200  
 actcactgag acaaaacaaa gaagagccaa agttccagag ggacctgaga g 1251

<210> 51  
 <211> 402  
 <212> DNA

<213> Homo sapien

<400> 51

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cgagcggccg cccgggcagg taccgcgtca gagattatcc acagcagcca gatgggttcta      60
ccttccacaa agattgtggt tgcaattctg ggcttctaag ttctggttac ttcataatctt      120
tccttttggt cctccagccc tagagggtgt agctgctttc tgaagttatt atttctagat      180
gacttttggt ttttcagcct ttgtatcttg cttttcagcc ctctaagcc tgtataacca      240
atttcctgt aactaaataa atttcctcca ttgaaaaaaa aaaaaaaaaa aaaaaaaaaa      300
ggttgtgtgg ggttattcgg tggctctagg gcgtgttccc tgtgtgtgtg gaatgtgggt      360
ttcccggtcc aaaatttccc caaaaaattg cggacacacc tg                          402
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<210> 52

<211> 1042

<212> DNA

<213> Homo sapien

<400> 52

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caattgttct caaacttcac tagccccgtc ggcgcgagc cttgtcgaga atgcagattc      60
ctgggtactg ccagatacga attgagcata ccacaaaaaa gttctcattt tgtgtcctcc      120
catcccatte tcctcactaa ccaaaggcta ggaattatct gtgaatgtag gaccactgga      180
tttgagctct tcctctgaca ctgtggagag tttctaggaa tgaaacagat atatggcctt      240
gggtcccttt tttttttctt tttttttttt ttaatagaga cgagcatctc actatgttgc      300
ctagggtagt cttgaactcc tggcctcaag caatccccac ccgactccgc ctctcgaagt      360
gatgggatta caggcataaa ccaccacgcc tggccagaag gtgctttaac accaaatctg      420
aaaattgttc agaagagaaa cattgagcat gaacaccatc tgtgcgagtc atttacttat      480
tgccctcac ctctaaatct accttctgta ctcttcttcc ctgtaatgat ggggctagtt      540
gtcctcaaac tgtttctcag acttcttttt aagcttgctt cctgttcagt tctgccaata      600
ggggtcacta gagagagact gggaggcaga aggagagaat atgcttctctg ttttttctgt      660
tcttgtaaat gttgcttaca ggaccagcaa tgcttcttca cctagagaca cttctcccag      720
cagtggcagt gccacttcag cttctttcag cactactgga atcagcctca gtgattcccc      780
ctgtaccgc tcagagatta tccacagcag ccagatgggt ctaccttcca caaagattgt      840
ggttgcaatt ctgggcttct aagttctggt tacttcatat ttttcctttt gttcctccag      900
ccctagaggt ggtagctgct ttctgaagtt attatttcta gatgactttt ggtttttcag      960
cctttgtatt ttgcttttca gccctctaata gcctgtataa ccaatttccc tgtaataaat     1020
caatttcctc cattgaaaaa aa                                              1042
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<210> 53  
 <211> 240  
 <212> DNA  
 <213> Homo sapien

<220>  
 <221> misc\_feature  
 <222> (44)..(44)  
 <223> a, c, g or t

<400> 53  
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 ttgaacacaa gggtcagttc ttcaattcat gagcagtcag aacaggagat gcttaggaag 120  
 gaatcgtagc tggtagcctc tctccatgct catcccatc cccagtgcac ggataccgtt 180  
 ccctgaagtt taaaaacatg caccacactt ccggtaaagg ctggagccac agaggcacct 240

<210> 54  
 <211> 1590  
 <212> DNA  
 <213> Homo sapien

<400> 54  
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 caggcatctg cagtgcagcg aggttatgga aacccattg aagccagttc gtatgggctg 120  
 gacctggact gcggagctcc tggcacccca gaggtcatg tctgttttga cccctgtcag 180  
 aattacaccc tcttgatga acccttcga agcacagaga actcagcagg gtcccagggg 240  
 tgcgataaaa acatgagcgg ctggtaccgc tttgtagggg aaggaggagt aaggatgtcg 300  
 gagacctgtg tccaggtgca ccgatgccag acagacgctc ccatgtggct gaatgggacc 360  
 caccctgccc ttggggatgg catcaccaac cacactgcct gtgccattg gaggggcaac 420  
 tgctgtttct ggaaaacaga ggtgctgggtg aaggcctgcc caggcgggta ccatgtgtac 480  
 cggttggaag gcactccctg gtgtaatctg agatactgca cagaccatc cactgtggag 540  
 gacaagtgtg agaaggcctg ccgccccgag gaggagtgcc ttgccctcaa cagcacctgg 600  
 ggctgtttct gcagacagga cctcaatagt tctgatgtcc acagtttgca gcctcagcta 660  
 gactgtgggc ccaggagat caaggtgaag gtggacaaat gtttgctggg aggcctgggt 720  
 ttgggggagg aggtcattgc ctacctgca gacccaaact gcagcagcat cttgcagaca 780  
 gaggagagga actgggtatc tgtgaccagc cccgtccagg ctagtgcctg caggaacatt 840  
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atcatcagag acaccatcct caacatcaac ttccaatgtg cctaccact ggacatgaaa 960  
gtcagcctcc aagctgcctt gcagcccatt gtaagttccc tgaacgtcag tgtggacggg 1020  
aatggagagt tcattgtcag gatggccctc ttccaagacc agaactacac gaatccttac 1080  
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caaggggaca cctcccgtt taacctggtg ttgaggaact gctatgccac cccactgaa 1200  
gacaaggctg accttgtgaa gtatttcac atcagaaaca gctgctcaaa tcaacgtgat 1260  
tccaccatcc acgtggagga gaatgggcag tccctcgaaa gccggttctc agttcagatg 1320  
ttcatgtttg ctggacatta tgacctagtt ttcctgcatt gtgagattca tctctgtgat 1380  
tctcttaatg aacagtgcc aacctcttgc tcaagaagtc aagtcgcag tgaagtaccg 1440  
gccatcgacc tagcccggt tctagatttg gggcccatca ctcgagagg tgcacagtct 1500  
cccgggtgca tgaatggaac ccttagcact gcagggttcc tgggtggcctg gcctatggtc 1560  
ctcctgactg tccctcctggc ttggctgttc 1590

<210> 55  
<211> 467  
<212> DNA  
<213> Homo sapien

<400> 55  
gtcgcggccg aggtacttat ataagggtta tttttaagtc caggaatttt ctcaaggaaa 60  
attttaagct actacaggcc aggtgcagtg gctcacacct gtaatcccag cactttggaa 120  
ggccaagggg gggcggtatc cgtaaggcca ggagttaaag accagcctgg ccaacatggc 180  
gaaaccccgct ctccactaaa aatacaaaaa ttagctgagg gtggtgggtgc atgtctgtaa 240  
tcccagctac tcgggaggtg gaggttgagc tgagctgaga tcacattgct tcaactccagc 300  
ctgggcgaca gagtgagact gtttaaaaaa aaattttttt aagctactgc aataaatttg 360  
tttattcatc aaataaaata aatagcaagg attttcttct attggaaaaa atagatagca 420  
aggattttct tctagtggaa aaagtttctc ctgtttaacc tggcatt 467

<210> 56  
<211> 2970  
<212> DNA  
<213> Homo sapien

<400> 56  
atgtcgggaag aaacccgaca gagcaaattg gccgcagcga agaaaaagtt gagagaatat 60  
cagcagagga atagccctgg tgttcctaca ggagcgaaaa agaagaagaa aataaaaaat 120  
ggcagtaacc ctgagacaac cacttctggt ggttgccact cacctgagga tacacccaag 180

gacaatgctg	ctactctaca	accatctgat	gacaccgtgt	tacctggcgg	tgteccctcc	240
cctgggtgcca	gtctcactag	catggcgcca	tctcagaatc	atgatgctga	caatgtccct	300
aatctcatgg	atgaaaccaa	gactttctca	tcaaccgaga	gcctgcgaca	actctcccaa	360
cagctcaatg	gtcttggttg	tgagtctgcg	acatgtgtca	atggggaggg	ccctgcatcg	420
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agctatgtaa	caaacaaaca	actcaatatc	acgatagaga	aattgaaaca	acagaaccaa	540
gaaattacgg	atcagttgga	agaagaaaag	aaagaatgcc	acaaaaagca	gggagcccta	600
agggagcagt	tacaggttca	cattcagacc	atagggatcc	tcgtatcaga	gaaagctgag	660
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agagacgccc	tcaggctgga	gttatacaag	aacacccaaa	gcaatgagga	cctgaagcaa	900
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cttaacttgg	aagaattgca	aaagaagtta	gagatgacgg	aactcctgct	tcaacagttt	1020
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aaggaggtgc	tgcataatca	gctactgctg	cagacccagc	tcgtggacca	gctgcagcag	1920

caggaagctc agggcaaagc ggtggccgag atggcccgcc aagagttgca ggaaacccag 1980  
gagcgcctgg aagctgccac ccagcagaat cagcagctac gggcccagtt gagcctcatg 2040  
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ccccgggagc gccaggctt gggcagcaac ccctgcattc ctttttttta ccgggctgac 2940  
gagaatgatg aggtgaagat cactgtcatc 2970

<210> 57  
<211> 461  
<212> DNA  
<213> Homo sapien

<400> 57  
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cctgggaagg tgaggcaggc cagagtctgt gttctgtgtt gagtgtcaag ctatttgcta 120  
ggaaggctct caacaggcct tgggtgtgggc tctgccagag actgttctga acacttgctt 180  
gagatccgtg ccctgtaaaa tggatatgat gttttactga tgtctgtaat acatttgtaa 240  
acttccaata aaatttgaat aaaagaaaaa taacaaaaaa caacaaaaaa aagaaaaaag 300  
aagcgcgggg cggtactgca ggggccatac gctgggtgtcc cgtgggggtg acatgggtga 360  
gatccgggtc aaaattccac ccaaactata gcgagcaatc ggagcatagc gacagagaag 420  
agagagcgac acagagatgc agacgaccaa agaacaggaa g 461

<210> 58  
 <211> 1032  
 <212> DNA  
 <213> Homo sapien

<400> 58  
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 caaaactagc tttcctcaag gaaatataaa ggaggggaaa gtcacatagt gttaggaaaa 120  
 cattcctgtg ttttgaatac gatgaatcca taggatagag aaaaatctgc ttgttctatt 180  
 ctgagagttc tctgagatat cccctcactc tgcttgccat ttggccattg atattcaaca 240  
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Asp Ile Lys Ser Tyr Lys Asp Phe Arg Phe Ser Phe Thr Lys Lys Val  
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Met Gly Lys Lys Ala His Arg His Leu Gln Phe Thr Ser Phe Lys Phe  
1 5 10 15

Leu Lys Lys Thr Pro Gln Lys Lys Pro Phe Leu Pro Gly Lys Ala His  
20 25 30

Glu Ile Asn Tyr Arg Ile Glu Leu Tyr Asn Ser Thr Ser Thr Ser Leu  
35 40 45



Met Ser Val Phe Phe Cys Val Lys Thr Pro Asp Thr Lys Thr Thr His  
1 5 10 15

Lys Thr Asn Lys Arg Lys Glu Asn Val Ala Arg Ile Leu Val Ser Leu  
                   20                                  25                                  30

Thr Val Glu Asp Pro Asp Gln Ala Val Gln Asn Val Ala His Gly Thr  
                   35                                  40                                  45

Glu Arg Thr Gly Val Thr Thr Glu Ile Lys Phe Val Gly Leu Gly Val  
                   50                                  55                                  60

Val Ala Pro Ser Gly  
 65

<210> 71  
 <211> 59  
 <212> PRT  
 <213> Homo sapien

<400> 71

Met Leu Ala Asp Ile Gly Val Leu Ile His Met Lys Trp Ile Asp Thr  
   1                                  5                                  10                                  15

Ser Ser Arg His His Thr Ala Val Gln Ser Ile Gln Gly Arg Glu Ala  
                   20                                  25                                  30

Thr Ser Arg Leu Thr Thr Phe Leu Ala Gly Ser Gly Glu Leu Cys Pro  
                   35                                  40                                  45

Arg Lys Pro Thr Arg Arg Ser Gly Thr Glu Glu  
                   50                                  55

<210> 72  
 <211> 50  
 <212> PRT  
 <213> Homo sapien

<400> 72

Met Phe Cys Ser Glu Asn Thr Leu Pro Gln Asp Ile Leu Gln Leu Ser  
   1                                  5                                  10                                  15

Tyr Cys Ile Gln Leu Ser Ala Gln Val Leu Thr Asp Glu Thr Cys His  
                   20                                  25                                  30

Pro Tyr Ser Thr Pro Cys Ser Ala Leu Leu Asn Ser Asn Ala His Met  
                   35                                  40                                  45

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Ala Ala Asp Ser Gly Phe Ser Ile Arg Gly Phe Ile Ile Ser Arg Thr  
65 70 75 80

Ser Ser Trp Ile Arg Val Ser Trp Ile Ser Cys Tyr Ser Asp Leu Trp  
85 90 95

Ala Glu Thr Thr Asn Asp Gly Thr Pro Gln Ser Thr Ser Pro Thr Ser  
100 105 110

Ala Ile His Thr Leu Ala Pro Arg Arg His Asp Leu Glu Ala His Arg  
115 120 125

Leu Ser Gly Tyr His  
130

<210> 75

<211> 72

<212> PRT

<213> Homo sapien

<400> 75

Met Trp Ser Val Ser Pro Cys Ser Leu Pro Glu Gln Cys Leu Arg Phe  
1 5 10 15

Glu Trp Asp Pro Thr Phe Val Asn Glu Ile Tyr His Leu Pro Arg Gln  
20 25 30

Asn Asn Arg Phe Cys Pro Arg Cys Cys Asp Val Thr Met Val Ala Ile  
35 40 45

Thr Ala Ile Thr Tyr Asn Tyr Trp His Thr Tyr Asp Glu Ser Arg Thr  
50 55 60

Gly Pro Lys Cys Phe Leu Thr Met  
65 70

<210> 76

<211> 93

<212> PRT

<213> Homo sapien

<400> 76

Met Ser Leu Cys Cys Asp Gly Pro Phe Pro Ser Leu Phe Gly Tyr Pro  
1 5 10 15

Pro Leu Thr Ile Leu Ile His Val Leu Phe Gln Lys Val Ser Pro Ile  
20 25 30

Lys Trp His Leu Gly Thr Thr Met Ala Gly Ile Ala Leu Ala Met Asn

35

40

45

Ser Thr Val Val Thr Leu Ser His Ser Arg Ala Val His Phe Ile Met  
50 55 60

Asn Asp Leu Arg Ile Ser Pro Gly Lys Ser Pro Arg Gln Ala Leu Pro  
65 70 75 80

Leu Leu Leu Ala Leu Gln Cys Glu Val Ser Trp Glu Arg  
85 90

<210> 77

<211> 500

<212> PRT

<213> Homo sapien

<400> 77

Met Lys Cys Thr Ala Arg Glu Trp Leu Arg Val Thr Thr Val Leu Phe  
1 5 10 15

Met Ala Arg Ala Ile Pro Ala Met Val Val Pro Asn Ala Thr Leu Leu  
20 25 30

Glu Lys Leu Leu Glu Lys Tyr Met Asp Glu Asp Gly Glu Trp Trp Ile  
35 40 45

Ala Lys Gln Arg Gly Lys Arg Ala Ile Thr Asp Asn Asp Met Gln Ser  
50 55 60

Ile Leu Asp Leu His Asn Lys Leu Arg Ser Gln Val Tyr Pro Thr Ala  
65 70 75 80

Ser Asn Met Glu Tyr Met Thr Trp Asp Val Glu Leu Glu Arg Ser Ala  
85 90 95

Glu Ser Trp Ala Glu Ser Cys Leu Trp Glu His Gly Pro Ala Ser Leu  
100 105 110

Leu Pro Ser Ile Gly Gln Asn Leu Gly Ala His Trp Gly Arg Tyr Arg  
115 120 125

Pro Pro Thr Phe His Val Gln Ser Trp Tyr Asp Glu Val Lys Asp Phe  
130 135 140

Ser Tyr Pro Tyr Glu His Glu Cys Asn Pro Tyr Cys Pro Phe Arg Cys

145					150					155					160				
Ser	Gly	Pro	Val	Cys	Thr	His	Tyr	Thr	Gln	Val	Val	Trp	Ala	Thr	Ser				
				165					170					175					
Asn	Arg	Ile	Gly	Cys	Ala	Ile	Asn	Leu	Cys	His	Asn	Met	Asn	Ile	Trp				
			180					185					190						
Gly	Gln	Ile	Trp	Pro	Lys	Ala	Val	Tyr	Leu	Val	Cys	Asn	Tyr	Ser	Pro				
		195					200					205							
Lys	Gly	Asn	Trp	Trp	Gly	His	Ala	Pro	Tyr	Lys	His	Gly	Arg	Pro	Cys				
	210					215					220								
Ser	Ala	Cys	Pro	Pro	Ser	Phe	Gly	Gly	Gly	Cys	Arg	Glu	Asn	Leu	Cys				
225					230					235					240				
Tyr	Lys	Glu	Gly	Ser	Asp	Arg	Tyr	Tyr	Pro	Pro	Arg	Glu	Glu	Glu	Thr				
				245					250					255					
Asn	Glu	Ile	Glu	Arg	Gln	Gln	Ser	Gln	Val	His	Asp	Thr	His	Val	Arg				
			260					265					270						
Thr	Arg	Ser	Asp	Asp	Ser	Ser	Arg	Asn	Glu	Val	Ile	Ser	Ala	Gln	Gln				
		275					280					285							
Met	Ser	Gln	Ile	Val	Ser	Cys	Glu	Val	Arg	Leu	Arg	Asp	Gln	Cys	Lys				
	290					295					300								
Gly	Thr	Thr	Cys	Asn	Arg	Tyr	Glu	Cys	Pro	Ala	Gly	Cys	Leu	Asp	Ser				
305					310					315					320				
Lys	Ala	Lys	Val	Ile	Gly	Ser	Val	His	Tyr	Glu	Met	Gln	Ser	Ser	Ile				
				325					330					335					
Cys	Arg	Ala	Ala	Ile	His	Tyr	Gly	Ile	Ile	Asp	Asn	Asp	Gly	Gly	Trp				
			340					345					350						
Val	Asp	Ile	Thr	Arg	Gln	Gly	Arg	Lys	His	Tyr	Phe	Ile	Lys	Ser	Asn				
		355					360					365							
Arg	Asn	Gly	Ile	Gln	Thr	Ile	Gly	Lys	Tyr	Gln	Ser	Ala	Asn	Ser	Phe				
	370					375					380								

55

Thr Val Ser Lys Val Thr Val Gln Ala Val Thr Cys Glu Thr Thr Val  
385 390 395 400

Glu Gln Leu Cys Pro Phe His Lys Pro Ala Ser His Cys Pro Arg Val  
405 410 415

Tyr Cys Pro Arg Asn Cys Met Gln Ala Asn Pro His Tyr Ala Arg Val  
420 425 430

Ile Gly Thr Arg Val Tyr Ser Asp Leu Ser Ser Ile Cys Arg Ala Ala  
435 440 445

Val His Ala Gly Val Val Arg Asn His Gly Gly Tyr Val Asp Val Met  
450 455 460

Pro Val Asp Lys Arg Lys Thr Tyr Ile Ala Ser Phe Gln Asn Gly Ile  
465 470 475 480

Phe Ser Glu Ser Leu Gln Asn Pro Pro Gly Gly Lys Ala Phe Arg Val  
485 490 495

Phe Ala Val Val  
500

<210> 78  
<211> 51  
<212> PRT  
<213> Homo sapien

<400> 78

Met Val Thr Thr Gln Asn Leu Arg Leu Thr Ile Val Glu Val Arg Gly  
1 5 10 15

Gln Gly Ala Gly Arg Ala Gly Ser Phe Leu Ser Ser Ile Met Gly Ala  
20 25 30

Ala Gly Arg Ile Gln Phe Leu Ala Gly Leu Gly Arg Arg Ser Pro Val  
35 40 45

Pro Ala Ala  
50

<210> 79  
<211> 50  
<212> PRT  
<213> Homo sapien

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&lt;400&gt; 79

Met Val Phe Tyr Tyr Tyr Tyr Gly Phe Lys Lys Ser Asn Phe Ile  
 1 5 10 15

Ser Phe Cys Lys Glu Leu Ser Asn Ile Leu Tyr Arg Phe Cys Glu Arg  
 20 25 30

Thr Tyr Phe Leu Thr Val Ile Phe Ile Ser Phe Lys Ile Phe Val Ser  
 35 40 45

His Leu  
 50

&lt;210&gt; 80

&lt;211&gt; 229

&lt;212&gt; PRT

&lt;213&gt; Homo sapien

&lt;400&gt; 80

Met Ala Glu Glu Met Glu Ser Ser Leu Glu Ala Ser Phe Ser Ser Ser  
 1 5 10 15

Gly Ala Val Ser Gly Ala Ser Gly Phe Leu Pro Pro Ala Arg Ser Arg  
 20 25 30

Ile Phe Lys Ile Ile Val Ile Gly Asp Ser Asn Val Gly Lys Thr Cys  
 35 40 45

Leu Thr Tyr Arg Phe Cys Ala Gly Arg Phe Pro Asp Arg Thr Glu Ala  
 50 55 60

Thr Ile Gly Val Asp Phe Arg Glu Arg Ala Val Glu Ile Asp Gly Glu  
 65 70 75 80

Arg Ile Lys Ile Gln Leu Trp Asp Thr Ala Gly Gln Glu Arg Phe Arg  
 85 90 95

Lys Ser Met Val Gln His Tyr Tyr Arg Asn Val His Ala Val Val Phe  
 100 105 110

Val Tyr Asp Met Thr Asn Met Ala Ser Phe His Ser Leu Pro Ser Trp  
 115 120 125

Ile Glu Glu Cys Lys Gln His Leu Leu Ala Asn Asp Ile Pro Arg Ile

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140

Ala Asn Phe His Thr Thr Ala Gly His Gly Ser Leu Leu Thr Thr His  
20 25 30

Cys His Leu Arg Leu Cys Leu Cys Phe Ile Gln Arg Glu Arg Gly Gly  
                   35                                  40                                  45

Leu Lys Gly Met Leu Pro Gly Gly  
           50                                  55

<210> . 83  
 <211> 72  
 <212> PRT  
 <213> Homo sapien

<400> 83

Met Leu Ser Pro Phe Leu Leu Ile Asn Asn Leu Tyr Tyr Lys Lys Lys  
 1                                  5                                  10                                  15

Lys Lys Lys Lys Lys Arg Arg Gly Gly Asn Gln Gly Pro Ile Arg Gly  
                   20                                  25                                  30

Phe Pro Gly Gly Glu Trp Val Thr Arg Ser Gln Phe His Thr Phe Ala  
                   35                                  40                                  45

Arg Gln Gln Thr Gly Glu Glu Ala Gly Pro Arg Arg Glu Ala Arg Gln  
           50                                  55                                  60

Glu Gln Ala His Arg Glu Thr Glu  
 65                                  70

<210> 84  
 <211> 27  
 <212> PRT  
 <213> Homo sapien

<400> 84

Met His Val Glu Arg Arg Ser Val Met Asp Ala Trp Ser Arg Arg Gly  
 1                                  5                                  10                                  15

Ala Gly Lys Tyr Thr Asp Ile Lys Asp Gln Ile  
           20                                  25

<210> 85  
 <211> 292  
 <212> PRT  
 <213> Homo sapien

<400> 85

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Met 1	Asn	Arg	Phe	Gly 5	Thr	Arg	Leu	Val	Gly 10	Ala	Thr	Ala	Thr	Ser 15	Ser
Pro	Pro	Pro	Lys 20	Ala	Arg	Ser	Asn	Glu 25	Asn	Leu	Asp	Lys	Ile 30	Asp	Met
Ser	Leu	Asp 35	Asp	Ile	Ile	Lys	Leu 40	Asn	Arg	Lys	Glu	Gly 45	Lys	Lys	Gln
Asn	Phe 50	Pro	Arg	Leu	Asn	Arg 55	Arg	Leu	Leu	Gln	Gln 60	Ser	Gly	Ala	Gln
Gln 65	Phe	Arg	Met	Arg	Val 70	Arg	Trp	Gly	Ile	Gln 75	Gln	Asn	Ser	Gly	Phe 80
Gly	Lys	Thr	Ser	Leu 85	Asn	Arg	Arg	Gly	Arg 90	Val	Met	Pro	Gly	Lys 95	Arg
Arg	Pro	Asn	Gly 100	Val	Ile	Thr	Gly	Leu 105	Ala	Ala	Arg	Lys	Thr 110	Thr	Gly
Ile	Arg	Lys 115	Gly	Ile	Ser	Pro	Met 120	Asn	Arg	Pro	Pro	Leu	Ser	Asp	Lys
Asn	Ile 130	Glu	Gln	Tyr	Phe	Pro 135	Val	Leu	Lys	Arg	Lys 140	Ala	Asn	Leu	Leu
Arg 145	Gln	Asn	Glu	Gly	Gln 150	Arg	Lys	Pro	Val	Ala 155	Val	Leu	Lys	Arg	Pro 160
Ser	Gln	Leu	Ser	Arg 165	Lys	Asn	Asn	Ile	Pro 170	Ala	Asn	Phe	Thr	Arg 175	Ser
Gly	Asn	Lys	Leu 180	Asn	His	Gln	Lys	Asp 185	Thr	Arg	Gln	Ala	Thr 190	Phe	Leu
Phe	Arg	Arg 195	Gly	Leu	Lys	Val	Gln 200	Ala	Gln	Leu	Asn	Thr 205	Glu	Gln	Leu
Leu	Asp 210	Asp	Val	Val	Ala	Lys 215	Arg	Thr	Arg	Gln	Trp 220	Arg	Thr	Ser	Thr
Thr 225	Asn	Gly	Gly	Ile 230	Leu	Thr	Val	Ser	Ile	Asp 235	Asn	Pro	Gly	Ala	Val 240

His Met Phe Glu Asp Phe Ser Phe Pro Phe Ala Ile Phe Leu Phe Phe

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61

1                    5                    10                    15

Leu Arg Arg Arg Ser Ala Leu Thr Pro Arg Leu Glu Ala Ser Gly Ala  
                  20                    25                    30

Ile Leu Ala Tyr Cys Asn Leu His Pro Pro Gly Ser Ser Asp Ser Pro  
                  35                    40                    45

Ala Ser Ala Ser Gly Val Ala Gly Ile Thr Gly Ala Arg His His Val  
                  50                    55                    60

Arg Leu Ile Phe Val Phe Ser Val Glu Thr Gly Phe Cys Tyr Val Gly  
65                    70                    75                    80

Gln Ala Gly Leu Lys Leu Leu Thr Ser Ser Asp Pro Pro Ala Ser Ala  
                  85                    90                    95

Ser Gln Ser Val Arg Ile Thr Gly Val Ser His Arg Ala Arg Leu Lys  
                  100                    105                    110

Ile Phe Leu Asn Cys Asn Lys Tyr Ser Ala Phe Phe Glu Ser Leu Tyr  
                  115                    120                    125

Leu

<210> 89  
<211> 15  
<212> PRT  
<213> Homo sapien

<400> 89

Met Ala Thr Leu Ala Gly Tyr Phe Leu Ala Lys Phe Leu Leu Arg  
1                    5                    10                    15

<210> 90  
<211> 71  
<212> PRT  
<213> Homo sapien

<400> 90

Met Lys His Gly Ser Phe Tyr Phe Thr Val Ser Asn Leu Ile Ala Ser  
1                    5                    10                    15

His Leu Lys Ser Ala Lys Ile Glu Leu Pro Lys Lys Cys Tyr Met Pro  
                  20                    25                    30

Lys Gly Ala His Asn Tyr Leu Met Ala Lys Leu Ile Lys Leu Thr Ser  
 35 40 45

Pro Lys Ser Asp Ser Arg Asp Leu Leu Cys Pro Ser Leu Trp Cys Phe  
 50 55 60

Phe Ala Leu His Ile Cys Phe  
 65 70

<210> 91  
 <211> 35  
 <212> PRT  
 <213> Homo sapien

<400> 91

Met Leu Ala Arg Leu Leu Met Ile Lys Ser Leu Asp Pro His Thr  
 1 5 10 15

Arg Phe Ala Met Val Thr Leu Ser Arg Thr Glu Ile Pro Leu Val Leu  
 20 25 30

Tyr Lys Arg  
 35

<210> 92  
 <211> 48  
 <212> PRT  
 <213> Homo sapien

<400> 92

Met Phe Thr Ser Thr Thr Leu Asn Gln Leu Leu Ser Ile Leu Tyr Ile  
 1 5 10 15

Phe Tyr Ser Ile Phe Phe Ser Asn Phe Leu His Phe Pro Met Ser Leu  
 20 25 30

Lys Phe Ser Val Asn Val Asn Phe Lys Asn Cys Thr Val Trp Leu Phe  
 35 40 45

<210> 93  
 <211> 67  
 <212> PRT  
 <213> Homo sapien

<400> 93

[illegible]

Met Cys Met Ser Arg Phe Glu Ser Leu Gly Cys Arg Phe Val Leu Pro  
1 5 10 15

Trp Gln Arg Lys Arg Ser Leu Trp Gly Gly Glu Leu Phe Leu Val Ile  
20 25 30

Ser Gly Lys Arg His Ile Glu Thr Leu Tyr Glu Trp Gly Phe Cys Phe  
35 40 45

Lys Cys Trp Lys Ile Arg Ala Gly Ile Thr Cys Leu Gln Val Val Pro  
50 55 60

Ser Leu Val  
65

<210>	94
<211>	145
<212>	PRT
<213>	Homo sapien

<400> 94

Met Leu Pro Ala Gly Thr Leu Val Gly Ala Gly Leu Gly Val Pro His  
1 5 10 15

Pro	Gln	Thr	Pro	Cys	Phe	Leu	Gln	Gly	His	Trp	Trp	Val	Leu	Ala	Trp
			20					25					30		

Gly Phe Leu Thr His Lys His His Ala Ser Cys Arg Asp Val Asp Gly  
35 40 45

Arg Trp Pro Gly Arg Ser Ser His Thr Thr Ala Met Leu Pro Ala Gly  
50 55 60

Thr Leu Val Gly Ala Gly Leu Gly Leu Pro His Ile Gln Thr Pro Cys  
65 70 75 80

Phe Leu Gln Gly Arg Trp Cys Ala Leu Ala Trp Gly Phe Leu Thr Tyr  
85 90 95

Lys Pro His Ala Ser Tyr Arg Ala Arg Trp Trp Thr Ala Gly Pro Glu  
100 105 110

Ala Ser Ser His Thr Ile Ala Ile Leu Pro His Gly Thr Leu Ala Ala  
115 120 125





&lt;400&gt; 97

Met Leu Arg Arg Glu Ala Arg Leu Arg Arg Glu Tyr Leu Tyr Arg Lys  
1 5 10 15

Ala Arg Glu Glu Ala Gln Arg Ser Ala Gln Glu Arg Lys Glu Arg Leu  
20 25 30

Arg Arg Ala Leu Glu Glu Asn Arg Leu Ile Pro Thr Glu Leu Arg Arg  
35 40 45

Glu Ala Leu Ala Leu Gln Gly Ser Leu Glu Phe Asp Asp Ala Gly Gly  
50 55 60

Glu Gly Val Thr Ser His Val Asp Asp Glu Tyr Arg Trp Ala Gly Val  
65 70 75 80

Glu Asp Pro Lys Val Met Ile Thr Thr Ser Arg Asp Pro Ser Ser Arg  
85 90 95

Leu Lys Met Phe Ala Lys Glu Leu Lys Leu Val Phe Pro Gly Ala Gln  
100 105 110

Arg Met Asn Arg Gly Arg His Glu Val Gly Ala Leu Val Arg Ala Cys  
115 120 125

Lys Ala Asn Gly Val Thr Asp Leu Leu Val Val His Glu His Arg Gly  
130 135 140

Thr Pro Val Gly Leu Ile Val Ser His Leu Pro Phe Gly Pro Thr Ala  
145 150 155 160

Tyr Phe Thr Leu Cys Asn Val Val Met Arg His Asp Ile Pro Asp Leu  
165 170 175

Gly Thr Met Ser Glu Ala Lys Pro His Leu Ile Thr His Gly Phe Ser  
180 185 190

Ser Arg Leu Gly Lys Arg Val Ser Asp Ile Leu Arg Tyr Leu Phe Pro  
195 200 205

Val Pro Lys Asp Asp Ser His Arg Val Ile Thr Phe Ala Asn Gln Asp  
210 215 220

Asp Tyr Ile Ser Phe Arg His His Val Tyr Lys Lys Thr Asp His Arg

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His Arg Pro Val Leu Asp Thr Leu Ala Met Leu Thr Ala His Arg Ala  
50 55 60

Gly Lys Lys His Leu Ser Ser Lys Leu Gly Gly Arg Arg Asp Gly Glu  
65 70 75 80

Ala Thr Leu Glu Ile Ser Ala His His Ser Trp Cys Tyr Ala Phe Asn  
85 90 95

Ser Val Ser Leu Ser Pro Gln Ala Leu Gln Leu Phe Tyr Gly Lys Lys  
100 105 110

Gln Pro Gly Lys Glu Arg Lys Gln Asn Pro Lys His Gln Asn Glu Leu  
115 120 125

Arg Arg Glu Glu Thr Lys Ala Glu Ala Pro Leu Leu Thr Gln Thr Arg  
130 135 140

Leu Ile Thr Gln Ser Ala Leu His Arg Ala Pro His Tyr Asn Ser Cys  
145 150 155 160

Cys Arg Arg Lys Tyr Arg Tyr Gly Thr Gly Lys Pro Glu Val  
165 170

<210> 100

<211> 50

<212> PRT

<213> Homo sapien

<400> 100

Met Lys Tyr Pro Phe Ile Tyr Asn Tyr Phe Cys Leu Lys His Val Ser  
1 5 10 15

Leu Tyr Ile Lys Asn Arg Tyr Phe Cys Phe His Phe Leu Ile Lys Phe  
20 25 30

Cys Pro Tyr Phe Arg Ser Glu Lys Asn Gln Tyr Ser Asn Ile Lys Lys  
35 40 45

Gln Glu  
50

<210> 101

<211> 18

<212> PRT

<213> Homo sapien

<400> 101

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Met Glu Glu Ile Tyr Leu Val Thr Gly Lys Leu Val Ile Gln Ala Leu  
 1 5 10 15

Glu Gly

<210> 102  
 <211> 34  
 <212> PRT  
 <213> Homo sapien

<400> 102

Met Ser Ser Gln Asn Arg Arg Cys Leu Gly Arg Asn Arg Gly Trp Cys  
 1 5 10 15

Leu Phe Ser Met Leu Ile Pro Tyr Pro Ser Asp Arg Ile Pro Phe Pro  
 20 25 30

Glu Val

<210> 103  
 <211> 40  
 <212> PRT  
 <213> Homo sapien

<400> 103

Met Asn Lys Gln Ile Tyr Cys Ser Ser Leu Lys Lys Phe Phe Phe Lys  
 1 5 10 15

Gln Ser His Ser Val Ala Gln Ala Gly Val Lys Gln Cys Asp Leu Ser  
 20 25 30

Ser Leu Gln Pro Pro Pro Pro Glu  
 35 40

<210> 104  
 <211> 990  
 <212> PRT  
 <213> Homo sapien

<400> 104

Met Ser Glu Glu Thr Arg Gln Ser Lys Leu Ala Ala Ala Lys Lys Lys  
 1 5 10 15

Leu Arg Glu Tyr Gln Gln Arg Asn Ser Pro Gly Val Pro Thr Gly Ala

30

Asp Leu Ala Ser Arg Leu Gln Tyr Ser Arg Arg Arg Val Gly Glu Leu  
245 250 255

$\frac{d}{dt} \left( \frac{\partial L}{\partial \dot{x}} \right) = \frac{\partial L}{\partial x}$

Glu Arg Ala Leu Ser Ala Val Ser Thr Gln Gln Lys Lys Ala Asp Arg  
                   260                  265                  270

Tyr Asn Lys Glu Leu Thr Lys Glu Arg Asp Ala Leu Arg Leu Glu Leu  
                   275                  280                  285

Tyr Lys Asn Thr Gln Ser Asn Glu Asp Leu Lys Gln Glu Lys Ser Glu  
                   290                  295                  300

Leu Glu Glu Lys Leu Arg Val Leu Val Thr Glu Lys Ala Gly Met Gln  
                   305                  310                  315                  320

Leu Asn Leu Glu Glu Leu Gln Lys Lys Leu Glu Met Thr Glu Leu Leu  
                   325                  330                  335

Leu Gln Gln Phe Ser Ser Arg Cys Glu Ala Pro Asp Ala Asn Gln Gln  
                   340                  345                  350

Leu Gln Gln Ala Met Glu Glu Arg Ala Gln Leu Glu Ala His Leu Gly  
                   355                  360                  365

Gln Val Met Glu Ser Val Arg Gln Leu Gln Met Glu Arg Asp Lys Tyr  
                   370                  375                  380

Ala Glu Asn Leu Lys Gly Glu Ser Ala Met Trp Arg Gln Arg Met Gln  
                   385                  390                  395                  400

Gln Met Ser Glu Gln Val His Thr Leu Arg Glu Glu Lys Glu Cys Ser  
                   405                  410                  415

Met Ser Arg Val Gln Glu Leu Glu Thr Ser Leu Ala Glu Leu Arg Asn  
                   420                  425                  430

Gln Met Ala Glu Pro Pro Pro Pro Glu Pro Pro Ala Gly Pro Ser Glu  
                   435                  440                  445

Val Glu Gln Gln Leu Gln Ala Glu Ala Glu His Leu Arg Lys Glu Leu  
                   450                  455                  460

Glu Gly Leu Ala Gly Gln Leu Gln Ala Gln Val Gln Asp Asn Glu Gly  
                   465                  470                  475                  480

Leu Ser Arg Leu Asn Arg Glu Gln Glu Glu Arg Leu Leu Glu Leu Glu  
                   485                  490                  495

Arg Ala Ala Glu Leu Trp Gly Glu Gln Ala Glu Ala Arg Arg Gln Ile  
500 505 510

Leu Glu Thr Met Gln Asn Asp Arg Thr Thr Ile Ser Arg Ala Leu Ser  
515 520 525

Gln Asn Arg Glu Leu Lys Glu Gln Leu Ala Glu Leu Gln Ser Gly Phe  
530 535 540

Val Lys Leu Thr Asn Glu Asn Met Glu Ile Thr Ser Ala Leu Gln Ser  
545 550 555 560

Glu Gln His Val Lys Arg Glu Leu Gly Lys Lys Leu Gly Glu Leu Gln  
565 570 575

Glu Lys Leu Ser Glu Leu Lys Glu Thr Val Glu Leu Lys Ser Gln Glu  
580 585 590

Ala Gln Ser Leu Gln Gln Gln Arg Asp Gln Tyr Leu Gly His Leu Gln  
595 600 605

Gln Tyr Val Ala Ala Tyr Gln Gln Leu Thr Ser Glu Lys Glu Val Leu  
610 615 620

His Asn Gln Leu Leu Leu Gln Thr Gln Leu Val Asp Gln Leu Gln Gln  
625 630 635 640

Gln Glu Ala Gln Gly Lys Ala Val Ala Glu Met Ala Arg Gln Glu Leu  
645 650 655

Gln Glu Thr Gln Glu Arg Leu Glu Ala Ala Thr Gln Gln Asn Gln Gln  
660 665 670

Leu Arg Ala Gln Leu Ser Leu Met Ala His Pro Gly Glu Gly Asp Gly  
675 680 685

Leu Asp Arg Glu Glu Glu Glu Asp Glu Glu Glu Glu Glu Glu Glu Ala  
690 695 700

Val Ala Val Pro Gln Pro Met Pro Ser Ile Pro Glu Asp Leu Glu Ser  
705 710 715 720

Arg Glu Ala Met Val Ala Phe Phe Asn Ser Ala Val Ala Ser Ala Glu  
725 730 735

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Glu Glu Gln Ala Arg Leu Arg Gly Gln Leu Lys Glu Gln Arg Val Arg  
740 745 750

Cys Arg Arg Leu Ala His Leu Leu Ala Ser Ala Gln Lys Glu Pro Glu  
755 760 765

Ala Ala Ala Pro Ala Pro Gly Thr Gly Gly Asp Ser Val Cys Gly Glu  
770 775 780

Thr His Arg Ala Leu Gln Gly Ala Met Glu Lys Leu Gln Ser Arg Phe  
785 790 795 800

Met Glu Leu Met Gln Glu Lys Ala Asp Leu Lys Glu Arg Val Glu Glu  
805 810 815

Leu Glu His Arg Cys Ile Gln Leu Ser Gly Glu Thr Asp Thr Ile Gly  
820 825 830

Glu Tyr Ile Ala Leu Tyr Gln Ser Gln Arg Ala Val Leu Lys Glu Arg  
835 840 845

His Arg Glu Lys Glu Glu Tyr Ile Ser Arg Leu Ala Gln Asp Lys Glu  
850 855 860

Glu Met Lys Val Lys Leu Leu Glu Leu Gln Glu Leu Val Leu Arg Leu  
865 870 875 880

Val Gly Asp Arg Asn Glu Trp His Gly Arg Phe Leu Ala Ala Ala Gln  
885 890 895

Asn Pro Ala Asp Glu Pro Thr Ser Gly Ala Pro Ala Pro Gln Glu Leu  
900 905 910

Gly Ala Ala Asn Gln Gln Gly Asp Leu Cys Glu Val Ser Leu Ala Gly  
915 920 925

Ser Val Glu Pro Ala Gln Gly Glu Ala Arg Glu Gly Ser Pro Arg Asp  
930 935 940

Asn Pro Thr Ala Gln Gln Ile Met Gln Leu Leu Arg Glu Met Gln Asn  
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Pro Arg Glu Arg Pro Gly Leu Gly Ser Asn Pro Cys Ile Pro Phe Phe

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Phe Phe Phe Ile Phe Leu Leu Phe Lys Phe Tyr Trp Lys Phe Thr Asn  
20 25 30

Val Leu Gln Thr Ser Val Lys His His Ile His Phe Thr Gly His Gly  
35 40 45

Ser Gln Ala Ser Val Gln Asn Ser Leu Trp Gln Ser Pro His Gln Gly  
50 55 60

Leu Leu Gln Thr Phe Leu Thr Asn Ser Leu Thr Leu Asn Thr Glu His  
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Arg Leu Trp Pro Ala Ser Pro Ser Gln Ala Leu  
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20 25 30

Ala Trp Val Leu Gly Glu Glu Ala Pro Gly Gln Arg Pro Pro Ala Ser  
35 40 45

Leu Gln Glu Ala Trp Gln Leu Tyr Val Arg Lys Pro Arg Pro Ala Pro  
50 55 60

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Ser Thr Leu Leu Phe Leu Ser Leu Ser Ser Leu Lys Met Tyr Asn Lys  
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Ile Ser Phe Leu Ala Pro Arg Leu Ser Pro Pro Gln Asn Lys Lys Lys  
35 40 45

Lys Lys Lys Lys Lys Asn Pro Phe Phe Phe Phe Phe Phe Phe Phe Leu  
50 55 60

Phe Phe Phe Phe Phe Phe Phe Ala His Asn Lys Asn Leu Leu Gly Glu  
65 70 75 80

Arg Trp Leu Met Gly Gly Lys Ile Trp Ile Gln Glu Ser Ser Ile Leu  
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Ala Leu Ala Leu Ser Pro Asn Pro Pro Ser Leu Pro Glu Pro Arg Gly  
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Val Ser Pro Cys  
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Ile Met Arg Thr Asp Asn Leu Pro Trp Ser Gln Arg Pro Ser Leu Pro  
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Thr Lys Asp Lys Ala Gln Ser Phe Phe Phe Phe Phe Phe Phe Phe  
                   20                                  25                                  30

Arg Leu Ser Thr Leu Leu Ser Cys Pro Gln Ala Pro Arg Asn Ile Leu  
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Ser Pro His Leu Glu Thr Asp  
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Ala Ser Ala Gly Ala Ala Gly Ser Leu Thr Arg Ser Pro Ser Ser Asp  
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Phe Gln Gly Ala Ser Val Glu Lys Lys Met Ala Gln Val Leu His Val  
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Pro Ala Pro Phe Pro Gly Thr Pro Gly Pro Ala Ser Pro Pro Ala Phe  
                   35                                  40                                  45

Pro Ala Lys Asp Pro Asp Pro Pro Tyr Ser Val Glu Thr Pro Tyr Gly  
                   50                                  55                                  60

Tyr Arg Leu Asp Leu Asp Phe Leu Lys Tyr Val Asp Asp Ile Glu Lys  
 65                                  70                                  75                                  80

Gly His Thr Leu Arg Arg Val Ala Val Gln Arg Arg Pro Arg Leu Ser  
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Cys Ser Asn Ala Ser Gly Asp Ser Arg His Ser Ala Tyr Ser Tyr Cys  
 115 120 125

Gly Arg Gly Phe Tyr Pro Gln Tyr Gly Ala Leu Glu Thr Arg Gly Gly  
 130 135 140

Phe Asn Pro Arg Val Glu Arg Thr Leu Leu Asp Ala Arg Arg Arg Leu  
 145 150 155 160

Glu Asp Gln Ala Ala Thr Pro Thr Gly Leu Gly Ser Leu Thr Pro Ser  
 165 170 175

Ala Ala Gly Ser Thr Ala Ser Leu Val Gly Val Gly Leu Pro Pro Pro  
 180 185 190

Thr Pro Arg Ser Ser Gly Leu Ser Thr Pro Val Pro Pro Ser Ala Gly  
 195 200 205

His Leu Ala His Val Arg Glu Gln Met Ala Gly Ala Leu Arg Lys Leu  
 210 215 220

Arg Gln Leu Glu Glu Gln Val Lys Leu Ile Pro Val Leu Gln Val Lys  
 225 230 235 240

Leu Ser Val Leu Gln Glu Glu Lys Arg Gln Leu Thr Val Gln Leu Lys  
 245 250 255

Ser Gln Lys Phe Leu Gly His Pro Thr Ala Gly Arg Gly Arg Ser Glu  
 260 265 270

Leu Cys Leu Asp Leu Pro Asp Pro Pro Glu Asp Pro Val Ala Leu Glu  
 275 280 285

Thr Arg Ser Val Gly Thr Trp Val Arg Glu Arg Asp Leu Gly Met Pro  
 290 295 300

Asp Gly Glu Ala Ala Leu Ala Ala Lys Val Ala Val Leu Glu Thr Gln  
 305 310 315 320

Leu Lys Lys Ala Leu Gln Glu Leu Gln Ala Ala Gln Ala Arg Gln Ala  
 325 330 335

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Asp Pro Gln Pro Gln Ala Trp Pro Pro Pro Asp Ser Pro Val Arg Val  
340 345 350

Asp Thr Val Arg Val Val Glu Gly Pro Arg Glu Val Glu Val Val Ala  
355 360 365

Ser Thr Ala Ala Gly Ala Pro Ala Gln Arg Ala Gln Ser Leu Glu Pro  
370 375 380

Tyr Gly Thr Gly Leu Arg Ala Leu Ala Met Pro Gly Arg Pro Glu Ser  
385 390 395 400

Pro Pro Val Phe Arg Ser Gln Glu Val Val Glu Thr Met Cys Pro Val  
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Pro Ala Ala Ala Thr Ser Asn Val His Met Val Lys Lys Ile Ser Ile  
420 425 430

Thr Glu Arg Ser Cys Asp Gly Ala Ala Gly Leu Pro Glu Val Pro Ala  
435 440 445

Glu Ser Ser Ser Ser Pro Pro Gly Ser Glu Val Ala Ser Leu Thr Gln  
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Pro Glu Lys Ser Thr Gly Arg Val Pro Thr Gln Glu Pro Thr His Arg  
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Glu Pro Thr Arg Gln Ala Ala Ser Gln Glu Ser Glu Glu Ala Gly Gly  
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Thr Gly Gly Pro Pro Ala Gly Val Arg Ser Ile Met Lys Arg Lys Glu  
500 505 510

Glu Val Ala Asp Pro Thr Ala His Arg Arg Ser Leu Gln Phe Val Gly  
515 520 525

Val Asn Gly Gly Tyr Glu Ser Ser Ser Glu Asp Ser Ser Thr Ala Glu  
530 535 540

Asn Ile Ser Asp Asn Asp Ser Thr Glu Asn Glu Ala Pro Glu Pro Arg  
545 550 555 560

Glu Arg Val Pro Ser Val Ala Glu Ala Pro Gln Leu Arg Pro Ala Gly  
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Thr Ala Ala Ala Lys Thr Ser Arg Gln Glu Cys Gln Leu Ser Arg Glu  
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Ser Gln His Ile Pro Thr Ala Glu Gly Ala Ser Gly Ser Asn Thr Glu  
595 600 605

Glu Glu Ile Arg Met Glu Leu Ser Pro Asp Leu Ile Ser Ala Cys Leu  
610 615 620

Ala Leu Glu Lys Tyr Leu Asp Asn Pro Asn Ala Leu Thr Glu Arg Glu  
625 630 635 640

Leu Lys Val Ala Tyr Thr Thr Val Leu Gln Glu Trp Leu Arg Leu Ala  
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Cys Arg Ser Asp Ala His Pro Glu Leu Val Arg Arg His Leu Val Thr  
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Phe Arg Ala Met Ser Ala Arg Leu Leu Asp Tyr Val Val Asn Ile Ala  
675 680 685

Asp Ser Asn Gly Asn Thr Ala Leu His Tyr Ser Val Ser His Ala Asn  
690 695 700

Phe Pro Val Val Gln Gln Leu Leu Asp Ser Gly Val Cys Lys Val Asp  
705 710 715 720

Lys Gln Asn Arg Ala Gly Tyr Ser Pro Ile Met Leu Thr Ala Leu Ala  
725 730 735

Thr Leu Lys Thr Gln Asp Asp Ile Glu Thr Val Leu Gln Leu Phe Arg  
740 745 750

Leu Gly Asn Ile Asn Ala Lys Ala Ser Gln Ala Gly Gln Thr Ala Leu  
755 760 765

Met Leu Ala Val Ser His Gly Arg Val Asp Val Val Lys Ala Leu Leu  
770 775 780

Ala Cys Glu Ala Asp Val Asn Val Gln Asp Asp Asp Gly Ser Thr Ala  
785 790 795 800

Leu Met Cys Ala Cys Glu His Gly His Lys Glu Ile Ala Gly Leu Leu

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805

810

815

Leu Ala Val Pro Ser Cys Asp Ile Ser Leu Thr Asp Arg Asp Gly Ser  
820 825 830

Thr Ala Leu Met Val Ala Leu Asp Ala Gly Gln Ser Glu Ile Ala Ser  
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Met Leu Tyr Ser Arg Met Asn Ile Lys Cys Ser Phe Ala Pro Met Ser  
850 855 860

Asp Asp Glu Ser Pro Thr Ser Ser Ser Ala Glu Glu  
865 870 875

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